



SEQUENCE LISTING

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Davey, Keith

<120> Characterization of CYP2D6 Genotypes

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19

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<400> 240
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19

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<400> 241
ggccccctgca ctgtttc

17

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<220>
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<223> The residue at this position is linked to a Z28 quenching group.

<400> 242
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33

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 <220>
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 <220>
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 <400> 244
 cccagctgga tgagctgcta actgagcat 29

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 <400> 245
 atgacgtggc agaccaggat gacctgggav 30

 <210> 246
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 cgcgccgagg cggatgacct gggav 25

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 <400> 248
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 ccgttggggc gaaaggggag tca 23

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 acggacgcgg agttgggggt gggagav 27

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 <400> 251
 atgacgtggc agacctgggg gtgggagv 28

<210> 252
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 cgcacacccc acccccaaga cgcacacccc gccccaacgg tc 42

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 acggacgcgg agaggaaccc tggacat 28

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 aggagtagec acgtcgggtg aggatcttca tt 32

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 cgcgcgcgagg caggtagtcg gtgagat 27

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 <400> 260
 cccgcgcgca cccactgag cc 22

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 <400> 261
 acggacgcgg agttacagca cagggtgc 27

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<220>
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<220>
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<220>
 <221> misc_feature
 <222> (275)..(275)
 <223> The residue at this position is c or t.

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 gtgcctttcc attcatgaca cttcttgaat ctctggcaga accagccagc cgtgttcaca 120
 gtgtcaaatg aagggatgtc tttgattgct tccaggtggt cctcagcacc accggagggg 180
 gatgggtgat cagccgaatc tttgactcgg gctacccatg ggacatgggtg ttcattgacac 240
 gcttttcagaa catgttgaga aattccctcc caacnccaat tgtgacttgg ttgatggagc 300
 gaaagataaa caactggctc aatcatgcaa attacggctt aataccagaa gacaggtaaa 360
 tataatgtga ctgccaaggg ctttttaggaa gaaggagcct ctgcctgtcc agcagcctat 420
 acaagccagg cagtaccaca gcaacatggc tgaatgtgtg ggaacacttg atacaaattt 480
 gcttgataat aacagctaac tggtcttaag tactcagaaa gtgaaattat gtatttc 537

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<400> 264
 ctgggctggg agcagcctc 19

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cactcgctgg cctgtttcat gtc

23

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<400> 266
ctggaatccg gtgtcgaagt gg

22

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<400> 267
ctcggcccct gcactgtttc

20

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<400> 268
gaggcaagaa ggagtgtcag gg

22

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agtcctgtgg tgaggtgacg agg

23

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<400> 270
gccaccatgg tgtctttgct ttc

23

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<400> 271
accggattcc agctgggaaa tg

22

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<400> 272
accgggcacc tgtactcctc a

21

<210> 273
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<400> 273
gcatgagcta aggcacccag ac

22

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acggacgcgg agttacagca caggtgc

27

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27

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42

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41

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<400> 288
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25

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cgcgcgagg agacgcccet ttcg

24

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51

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51

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21

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28

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38

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38

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27

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32

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22

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20

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gaggcaagaa ggagtgtcag gg

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23

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15

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16

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19

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15

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12

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13

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14

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17

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17

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16

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17

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